SEQUENCE LISTING

<110> Rothman,James
Mayhew, Mark
Hoe, Mee

<120> KDEL RECEPTOR INHIBITORS

<130> 31488

<160> 36

<170> FastSEQ for Windows Version 3.0

<210> 1
<211> 46
<212> PRT
<213> rat

<400> 1

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Gly Asp Leu Ala Pro Gln Met Leu Arg Glu Leu Gln Glu Thr Asn Ala
 1               5                  10                  15
Ala Leu Gln Asp Val Arg Glu Leu Leu Arg Gln Gln Val Lys Glu Ile
            20                  25                  30
Thr Phe Leu Lys Asn Thr Val Met Glu Cys Asp Ala Cys Gly
        35                  40                  45
```

<210> 2
<211> 46
<212> PRT
<213> human

<400> 2

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Ser Asp Leu Gly Pro Gln Met Leu Arg Glu Leu Gln Glu Thr Asn Ala
 1               5                  10                  15
Ala Leu Gln Asp Val Arg Asp Trp Leu Arg Gln Gln Val Arg Glu Ile
            20                  25                  30
Thr Phe Leu Lys Asn Thr Val Met Glu Cys Asp Ala Cys Gly
        35                  40                  45
```

<210> 3
<211> 46
<212> PRT
<213> mouse

<400> 3

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Gly Glu Gln Thr Lys Ala Leu Val Thr Gln Leu Thr Leu Phe Asn Gln
 1               5                  10                  15
Ile Leu Val Glu Leu Arg Asp Asp Ile Arg Asp Gln Val Lys Glu Met
```

```
            20                  25                  30
Ser Leu Ile Arg Asn Thr Ile Met Glu Cys Gln Val Cys Gly
        35                  40                  45

      <210> 4
      <211> 46
      <212> PRT
      <213> human

      <400> 4
Gly Glu Gln Thr Lys Ala Leu Val Thr Gln Leu Thr Leu Phe Asn Gln
 1               5                  10                  15
Ile Leu Val Glu Leu Arg Asp Asp Ile Arg Asp Gln Val Lys Glu Met
            20                  25                  30
Ser Leu Ile Arg Asn Thr Ile Met Glu Cys Gln Val Cys Gly
        35                  40                  45

      <210> 5
      <211> 46
      <212> PRT
      <213> human

      <400> 5
Gly Asp Phe Asn Arg Gln Phe Leu Gly Gln Met Thr Gln Leu Asn Gln
 1               5                  10                  15
Leu Leu Gly Glu Val Lys Asp Leu Leu Arg Gln Gln Val Lys Glu Thr
            20                  25                  30
Ser Phe Leu Arg Asn Thr Ile Ala Glu Cys Gln Ala Cys Gly
        35                  40                  45

      <210> 6
      <211> 46
      <212> PRT
      <213> xenopus laevis

      <400> 6
Gly Asp Val Ser Arg Gln Leu Ile Gly Gln Ile Thr Gln Met Asn Gln
 1               5                  10                  15
Met Leu Gly Glu Leu Arg Asp Val Met Arg Gln Gln Val Lys Glu Thr
            20                  25                  30
Met Phe Leu Arg Asn Thr Ile Ala Glu Cys Gln Ala Cys Gly
        35                  40                  45

      <210> 7
      <211> 27
      <212> PRT
      <213> human

      <400> 7
Gln Lys Leu Gln Asn Leu Phe Ile Asn Phe Cys Leu Ile Leu Ile Cys
 1               5                  10                  15
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Leu Leu Leu Ile Cys Ile Ile Val Met Leu Leu
20 25

<210> 8
<211> 9
<212> PRT
<213> human papilloma virus

<400> 8
Leu Leu Leu Gly Thr Leu Asn Ile Val
1 5

<210> 9
<211> 9
<212> PRT
<213> human papilloma virus

<400> 9
Leu Leu Met Gly Thr Leu Gly Ile Val
1 5

<210> 10
<211> 9
<212> PRT
<213> human papilloma virus

<400> 10
Thr Leu Gln Asp Ile Val Leu His Leu
1 5

<210> 11
<211> 9
<212> PRT
<213> human papilloma virus

<400> 11
Gly Leu His Cys Tyr Glu Gln Leu Val
1 5

<210> 12
<211> 9
<212> PRT
<213> human papilloma virus

<400> 12
Pro Leu Lys Gln His Phe Gln Ile Val
1 5

<210> 13
<211> 115
<212> PRT

<213> Artificial Sequence

<220>
<223> chimeric rat comp

<400> 13

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Met Gly Lys Phe Thr Val Val Ala Ala Ala Leu Leu Leu Leu Gly Ala
 1               5                  10                  15
Val Arg Ala Glu Gly Ser Ser Leu Gly Gly Asp Leu Ala Pro Gln Met
            20                  25                  30
Leu Arg Glu Leu Gln Glu Thr Asn Ala Ala Leu Gln Asp Val Arg Glu
        35                  40                  45
Leu Leu Arg Gln Gln Val Lys Glu Ile Thr Phe Leu Lys Asn Thr Val
    50                  55                  60
Met Glu Cys Asp Ala Cys Gly Met Gln Pro Ala Arg Thr Pro Gly Thr
65                  70                  75                  80
Ser Pro Gln Pro Gln Pro Lys Pro Gln Pro Gln Pro Gln Pro Gln Pro
                85                  90                  95
Lys Pro Gln Pro Lys Pro Glu Pro Glu Gly Thr Gly Ser Ser Glu Lys
            100                 105                 110
Asp Glu Leu
        115
```

<210> 14
<211> 387
<212> DNA
<213> Artificial Sequence

<220>
<223> chimeric rat COMP-KDEL

<400> 14

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aagcttacca tgggaaagtt cactgtggtg gcggcggcgt tgctgctgct gggcgcggtg      60
cgggccgagg gatccagcct gggtggagac ctagccccac agatgcttcg agaactccag     120
gagactaatg cggcgctgca agacgtgaga gagctcttgc gacagcaggt caaggagatc     180
accttcctga agaatacggt gatggaatgt gacgcttgcg gaatgcagcc cgcacgcacc     240
cccggtacta gtccgcagcc gcagccgaaa ccgcagccgc agccgcagcc gcagccgaaa     300
ccgcagccga aaccggaacc ggaaggtacc ggatcatcag aaaaagatga gttgtaggcg     360
gccgcagaat tccatatgca tctcgag                                        387
```

<210> 15
<211> 115
<212> PRT
<213> Artificial Sequence

<220>
<223> chimeric rat COMP-KDEL

<400> 15

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Met Gly Lys Phe Thr Val Val Ala Ala Ala Leu Leu Leu Leu Gly Ala
 1               5                  10                  15
```

```
Val Arg Ala Glu Gly Ser Ser Leu Gly Gly Asp Cys Cys Pro Gln Met
            20                  25                  30
Leu Arg Glu Leu Gln Glu Thr Asn Ala Ala Leu Gln Asp Val Arg Glu
        35                  40                  45
Leu Leu Arg Gln Gln Val Lys Glu Ile Thr Phe Leu Lys Asn Thr Val
    50                  55                  60
Met Glu Cys Asp Ala Cys Gly Met Gln Pro Ala Arg Thr Pro Gly Thr
65                  70                  75                  80
Ser Pro Gln Pro Gln Pro Lys Pro Gln Pro Gln Pro Gln Pro Gln Pro
                85                  90                  95
Lys Pro Gln Pro Lys Pro Glu Pro Glu Gly Thr Gly Ser Ser Glu Lys
            100                 105                 110
Asp Glu Leu
        115
```

<210> 16
<211> 387
<212> DNA
<213> Artificial Sequence

<220>
<223> chimeric rat COMP-KDEL

<400> 16

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aagcttacca tgggaaagtt cactgtggtg gcggcggcgt tgctgctgct gggcgcggtg      60
cgggccgagg gatccagcct gggtggagac tgttgtccac agatgcttcg agaactccag     120
gagactaatg cggcgctgca agacgtgaga gagctcttgc gacagcaggt caaggagatc     180
accttcctga agaatacggt gatggaatgt gacgcttgcg gaatgcagcc cgcacgcacc     240
cccggtacta gtccgcagcc gcagccgaaa ccgcagccgc agccgcagcc gcagccgaaa     300
ccgcagccga aaccggaacc ggaaggtacc ggatcatcag aaaaagatga gttgtaggcg     360
gccgcagaat tccatatgca tctcgag                                        387
```

<210> 17
<211> 105
<212> PRT
<213> Artificial Sequence

<220>
<223> chimeric mouse TSP3-KDEL

<400> 17

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Met Gly Lys Phe Thr Val Val Ala Ala Ala Leu Leu Leu Leu Gly Ala
 1               5                  10                  15
Val Arg Ala Glu Gly Ser Ser Leu Gly Gly Asp Cys Cys Lys Ala Leu
            20                  25                  30
Val Thr Gln Leu Thr Leu Phe Asn Gln Ile Leu Val Glu Leu Arg Asp
        35                  40                  45
Asp Ile Arg Asp Gln Val Lys Glu Met Ser Leu Ile Arg Asn Thr Ile
    50                  55                  60
Met Glu Cys Gln Val Cys Gly Pro Gln Pro Gln Pro Lys Pro Gln Pro
65                  70                  75                  80
```

Gln Pro Gln Pro Gln Pro Lys Pro Gln Pro Lys Pro Glu Pro Glu Gly
85 90 95
Thr Gly Ser Ser Glu Lys Asp Glu Leu
100 105

<210> 18
<211> 357
<212> DNA
<213> Artificial Sequence

<220>
<223> chimeric mouse TSP3-KDEL

<400> 18
aagcttacca tgggaaagtt cactgtggtg gcggcggcgt tgctgctgct gggcgcggtg 60
cgggccgagg gatccagcct gggtggagac tgttgtaagg cattggtcac ccagctcacc 120
ctcttcaacc agatcctagt ggagcttcgg gacgacatcc gagaccaggt gaaggaaatg 180
tcactcatcc ggaacaccat catggagtgt caggtgtgcg gtccgcagcc gcagccgaaa 240
ccgcagccgc agccgcagcc gcagccgaaa ccgcagccga aaccggaacc ggaaggtacc 300
ggatcatcag aaaaagatga gttgtaggcg gccgcagaat tccatatgca tctcgag 357

<210> 19
<211> 109
<212> PRT
<213> Artificial Sequence

<220>
<223> chimeric mouse TSP3-KDEL

<400> 19
Met Gly Lys Phe Thr Val Val Ala Ala Ala Leu Leu Leu Leu Gly Ala
1 5 10 15
Val Arg Ala Glu Gly Ser Ser Leu Gly Gly Asp Cys Cys Gly Glu Gln
20 25 30
Thr Lys Ala Leu Val Thr Gln Leu Thr Leu Phe Asn Gln Ile Leu Val
35 40 45
Glu Leu Arg Asp Asp Ile Arg Asp Gln Val Lys Glu Met Ser Leu Ile
50 55 60
Arg Asn Thr Ile Met Glu Cys Gln Val Cys Gly Pro Gln Pro Gln Pro
65 70 75 80
Lys Pro Gln Pro Gln Pro Gln Pro Gln Pro Lys Pro Gln Pro Lys Pro
85 90 95
Glu Pro Glu Gly Thr Gly Ser Ser Glu Lys Asp Glu Leu
100 105

<210> 20
<211> 369
<212> DNA
<213> Artificial Sequence

<220>

<223> chimeric mouse TSP3-KDEL

<400> 20

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aagcttacca tgggaaagtt cactgtggtg gcggcggcgt tgctgctgct gggcgcggtg      60
cgggccgagg gatccagcct gggtggagac tgttgtgggg agcagaccaa ggcattggtc     120
acccagctca ccctcttcaa ccagatccta gtggagcttc gggacgacat ccgagaccag     180
gtgaaggaaa tgtcactcat ccggaacacc atcatggagt gtcaggtgtg cggtccgcag     240
ccgcagccga aaccgcagcc gcagccgcag ccgcagccga aaccgcagcc gaaaccggaa     300
ccggaaggta ccggatcatc agaaaaagat gagttgtagg cggccgcaga attccatatg     360
catctcgag                                                             369
```

<210> 21
<211> 109
<212> PRT
<213> Artificial Sequence

<220>
<223> chimeric Xenopus laevis TSP4-KDEL

<400> 21

```
Met Gly Lys Phe Thr Val Val Ala Ala Ala Leu Leu Leu Leu Gly Ala
 1               5                  10                  15
Val Arg Ala Glu Gly Ser Ser Leu Gly Gly Asp Cys Cys Gly Asp Val
            20                  25                  30
Ser Arg Gln Leu Ile Gly Gln Ile Thr Gln Met Asn Gln Met Leu Gly
        35                  40                  45
Glu Leu Arg Asp Val Met Arg Gln Gln Val Lys Glu Thr Met Phe Leu
    50                  55                  60
Arg Asn Thr Ile Ala Glu Cys Gln Ala Cys Gly Pro Gln Pro Gln Pro
65                  70                  75                  80
Lys Pro Gln Pro Gln Pro Gln Pro Gln Pro Lys Pro Gln Pro Lys Pro
                85                  90                  95
Glu Pro Glu Gly Thr Gly Ser Ser Glu Lys Asp Glu Leu
            100                 105
```

<210> 22
<211> 369
<212> DNA
<213> Artificial Sequence

<220>
<223> chimeric Xenopus laevis TSP4-KDEL

<400> 22

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aagcttacca tgggaaagtt cactgtggtg gcggcggcgt tgctgctgct gggcgcggtg      60
cgggccgagg gatccagcct gggtggagac tgttgtggtg acgtcagcag acagttgatt     120
ggccagataa cccaaatgaa tcagatgctg ggagagctcc gagatgtcat gagacagcag     180
gtgaaagaga ccatgttctt gagaaacacc attgcagaat gccaggcctg tggcccgcag     240
ccgcagccga aaccgcagcc gcagccgcag ccgcagccga aaccgcagcc gaaaccggaa     300
ccggaaggta ccggatcatc agaaaaagat gagttgtagg cggccgcaga attccatatg     360
catctcgag                                                             369
```

<210> 23
<211> 109
<212> PRT
<213> Artificial Sequence

<220>
<223> chimeric human COMP-KDEL

<400> 23

```
Met Arg Tyr Met Ile Leu Gly Leu Leu Ala Leu Ala Ala Val Cys Ser
 1               5                  10                  15
Ala Ala Lys Lys Gly Ser Ser Leu Gly Gly Asp Cys Cys Ser Asp Leu
            20                  25                  30
Gly Pro Gln Met Leu Arg Glu Leu Gln Glu Thr Asn Ala Ala Leu Gln
        35                  40                  45
Asp Val Arg Asp Trp Leu Arg Gln Gln Val Arg Glu Ile Thr Phe Leu
    50                  55                  60
Lys Asn Thr Val Met Glu Cys Asp Ala Cys Gly Pro Gln Pro Gln Pro
65                  70                  75                  80
Lys Pro Gln Pro Gln Pro Gln Pro Gln Pro Lys Pro Gln Pro Lys Pro
                85                  90                  95
Glu Pro Glu Gly Thr Gly Ser Ser Glu Lys Asp Glu Leu
            100                 105
```

<210> 24
<211> 372
<212> DNA
<213> Artificial Sequence

<220>
<223> chimeric human COMP-KDEL

<400> 24

```
aagcttacca tgggaaggta catgatttta ggcttgctcg cccttgcggc agtctgcagc      60
gctgccaaaa aaggatccag cctgggtgga gactgttgtt cagacctggg cccgcagatg     120
cttcgggaac tgcaggaaac caacgcggcg ctgcaggacg tgcgggactg gctgcggcag     180
caggtcaggg agatcacgtt cctgaaaaac acggtgatgg agtgtgacgc gtgcgggccg     240
cagccgcagc cgaaaccgca gccgcagccg cagccgcagc cgaaaccgca gccgaaaccg     300
gaaccggaag gtaccggatc atcagaaaaa gatgagttgt aggcggccgc agaattccat     360
atgcatctcg ag                                                        372
```

<210> 25
<211> 90
<212> PRT
<213> Artificial Sequence

<220>
<223> chimeric human PLB-KDEL

<400> 25

```
Met Arg Tyr Met Ile Leu Gly Leu Leu Ala Leu Ala Ala Val Cys Ser
```

```
 1               5                  10                  15
Ala Ala Lys Lys Gly Ser Ser Leu Gly Gly Asp Cys Cys Gln Lys Leu
            20                  25                  30
Gln Asn Leu Phe Ile Asn Phe Cys Leu Ile Leu Ile Cys Leu Leu Leu
        35                  40                  45
Ile Cys Ile Ile Val Met Leu Leu Pro Gln Pro Gln Pro Lys Pro Gln
    50                  55                  60
Pro Gln Pro Gln Pro Gln Pro Lys Pro Gln Pro Lys Pro Glu Pro Glu
65                  70                  75                  80
Gly Thr Gly Ser Ser Glu Lys Asp Glu Leu
                85                  90
```

<210> 26
<211> 315
<212> DNA
<213> Artificial Sequence

<220>
<223> chimeric human PLB-KDEL

<400> 26

```
aagcttacca tgggaaggta catgatttta ggcttgctcg cccttgcggc agtctgcagc      60
gctgccaaaa aaggatccag cctgggtgga gactgttgtc aaaagctaca gaatctattt     120
atcaatttct gtctcatctt aatatgtctc ttgctgatct gtatcatcgt gatgcttctc     180
ccgcagccgc agccgaaacc gcagccgcag ccgcagccgc agccgaaacc gcagccgaaa     240
ccggaaccgg aaggtaccgg atcatcagaa aaagatgagt tgtaggcggc cgcagaattc     300
catatgcatc tcgag                                                     315
```

<210> 27
<211> 109
<212> PRT
<213> Artificial Sequence

<220>
<223> chimeric human TSP3-KDEL

<400> 27

```
Met Arg Tyr Met Ile Leu Gly Leu Leu Ala Leu Ala Ala Val Cys Ser
 1               5                  10                  15
Ala Ala Lys Lys Gly Ser Ser Leu Gly Gly Asp Cys Cys Gly Glu Gln
            20                  25                  30
Thr Lys Ala Leu Val Thr Gln Leu Thr Leu Phe Asn Gln Ile Leu Val
        35                  40                  45
Glu Leu Arg Asp Asp Ile Arg Asp Gln Val Lys Glu Met Ser Leu Ile
    50                  55                  60
Arg Asn Thr Ile Met Glu Cys Gln Val Cys Gly Pro Gln Pro Gln Pro
65                  70                  75                  80
Lys Pro Gln Pro Gln Pro Gln Pro Gln Pro Lys Pro Gln Pro Lys Pro
                85                  90                  95
Glu Pro Glu Gly Thr Gly Ser Ser Glu Lys Asp Glu Leu
            100                 105
```

<210> 28
<211> 372
<212> DNA
<213> Artificial Sequence

<220>
<223> chimeric human TSP3-KDEL

<400> 28
aagcttacca tgggaaggta catgatttta ggcttgctcg cccttgcggc agtctgcagc 60
gctgccaaaa aaggatccag cctgggtgga gactgttgtg gggagcagac caaggcattg 120
gtcacccagc tcaccctctt caaccagatc ctagtggagc ttcgggacga catccgagac 180
caggtgaagg aaatgtcact catccggaac accatcatgg agtgtcaggt gtgcggtccg 240
cagccgcagc cgaaaccgca gccgcagccg cagccgcagc cgaaaccgca gccgaaaccg 300
gaaccggaag gtaccggatc atcagaaaaa gatgagttgt aggcggccgc agaattccat 360
atgcatctcg ag 372

<210> 29
<211> 109
<212> PRT
<213> Artificial Sequence

<220>
<223> chimeric human TSP4-KDEL

<400> 29
Met Arg Tyr Met Ile Leu Gly Leu Leu Ala Leu Ala Ala Val Cys Ser
1 5 10 15
Ala Ala Lys Lys Gly Ser Ser Leu Gly Gly Asp Cys Cys Gly Asp Phe
20 25 30
Asn Arg Gln Phe Leu Gly Gln Met Thr Gln Leu Asn Gln Leu Leu Gly
35 40 45
Glu Val Lys Asp Leu Leu Arg Gln Gln Val Lys Glu Thr Ser Phe Leu
50 55 60
Arg Asn Thr Ile Ala Glu Cys Gln Ala Cys Gly Pro Gln Pro Gln Pro
65 70 75 80
Lys Pro Gln Pro Gln Pro Gln Pro Gln Pro Lys Pro Gln Pro Lys Pro
85 90 95
Glu Pro Glu Gly Thr Gly Ser Ser Glu Lys Asp Glu Leu
100 105

<210> 30
<211> 372
<212> DNA
<213> Artificial Sequence

<220>
<223> chimeric human TSP4-KDEL

<400> 30
aagcttacca tgggaaggta catgatttta ggcttgctcg cccttgcggc agtctgcagc 60

gctgccaaaa aaggatccag cctgggtgga gactgttgtg gggactttaa ccggcagttc 120
ttgggtcaaa tgacacaatt aaaccaactc ctgggagagg tgaaggacct tctgagacag 180
caggttaagg aaacatcatt tttgcgaaac accatagctg aatgccaggc ttgcggtccg 240
cagccgcagc cgaaaccgca gccgcagccg cagccgcagc cgaaaccgca gccgaaaccg 300
gaaccggaag gtaccggatc atcagaaaaa gatgagttgt aggcggccgc agaattccat 360
atgcatctcg ag 372

<210> 31
<211> 8
<212> PRT
<213> unknown

<400> 31
Tyr Thr Ser Glu Lys Asp Glu Leu
1 5

<210> 32
<211> 8
<212> PRT
<213> unknown

<400> 32
Leu Asn Tyr Phe Asp Asp Glu Leu
1 5

<210> 33
<211> 9
<212> PRT
<213> unknown

<400> 33
Cys Asp Cys Arg Gly Asp Cys Phe Cys
1 5

<210> 34
<211> 134
<212> PRT
<213> Artificial Sequence

<220>
<223> KDEL/myc

<400> 34
Met Gly Lys Phe Thr Val Val Ala Ala Ala Leu Leu Leu Leu Gly Ala
1 5 10 15
Val Arg Ala Glu Gly Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
20 25 30
Tyr His Pro Asn Ser Thr Cys Gly Ser Ser Leu Gly Gly Asp Cys Cys
35 40 45
Pro Gln Met Leu Arg Glu Leu Gln Glu Thr Asn Ala Ala Leu Gln Asp
50 55 60

```
Val Arg Glu Leu Leu Arg Gln Gln Val Lys Glu Ile Thr Phe Leu Lys
65                  70                  75                  80
Asn Thr Val Met Glu Cys Asp Ala Cys Gly Met Gln Pro Ala Arg Thr
                85                  90                  95
Pro Gly Thr Ser Pro Gln Pro Gln Pro Lys Pro Gln Pro Gln Pro Gln
            100                 105                 110
Pro Gln Pro Lys Pro Gln Pro Lys Pro Glu Pro Glu Gly Thr Gly Ser
        115                 120                 125
Ser Glu Lys Asp Glu Leu
    130
```

<210> 35
<211> 444
<212> DNA
<213> Artificial Sequence

<220>
<223> KDEL-myc

<400> 35

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aagcttacca tgggaaagtt cactgtggtg gcggcggcgt tgctgctgct gggcgcggtg      60
cgggccgagg gatccgaaca aaaacttatt tctgaagaag acttgtacca cccaaactca     120
acatgcggat ccagcctggg tggagactgt tgtccacaga tgcttcgaga actccaggag     180
actaatgcgg cgctgcaaga cgtgagagag ctcttgcgac agcaggtcaa ggagatcacc     240
ttcctgaaga atacggtgat ggaatgtgac gcttgcggaa tgcagcccgc acgcaccccc     300
ggtactagtc cgcagccgca gccgaaaccg cagccgcagc cgcagccgca gccgaaaccg     360
cagccgaaac cggaaccgga aggtaccgga tcatcagaaa aagatgagtt gtaggcggcc     420
gcagaattcc atatgcatct cgag                                           444
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<210> 36
<211> 10
<212> PRT
<213> human myc tag

<400> 36

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Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
 1               5                  10
```